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GenCore version 5.1.3
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OM nucleic - nucleic search, using sw model

December 6, 2002, 22:41:46; Search time 2691.5 Seconds (without alignments) 16489.608 Million cell updates/sec Run on:

1 tctagaccatgtctggaaag.......ccaactcagaagtagtcgac 1525 US-10-025-514-7 1525 Title: Perfect score:

IDENTITY\_NUC Gapop 10.0 , Gapext 1.0 Scoring table: Sequence:

2054640 seqs, 14551402878 residues Searched:

4109280 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

GenEmb1:\*

Database :

9b\_bb: \*
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em\_or:\*
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em\_htg\_rvi:\*
em\_sy:\*
em\_sy:\*
em\_htgo\_rtc:\*

em\_htgo\_mus:\* em\_htgo\_other:\*

Pred. No. is the number of results predicted by chance to have a

score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

	Description	412 S	AR111411 Sequence	M11465 Human alpha	E00631 DNA encodin	edneuc		AX335339 Sequence	₽,	AX335338 Sequence	X01683 Human mRNA	AK026174 Homo sap1	E00195 cDNA encod1	I04196 Sequence 3	IO4272 Sequence 1	IO7849 Sequence 2	Sedne	9 ношо	uman Z	edneuc	HOMO S					I03509 Sequence 6		ednence 1	J00321 Papio hamad	I01227 Sequence 2	A01846 Human mRNA	ABOU4044 Cercopith	AXZUZUBY Sequence	בלתבווכם ד	edneuce	r admente	٠,		v i	S. Scrota m	7 Rat alp	2 alpha-1-an	75 Rat mRNA	B000546 Tamias	Tamias as	3129 B.taurus m
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## ALIGNMENTS

Sequence 6 from patent US 6127145. AR111412.1 GI:12828260 Unknown. RESULT 1
ARI11412
LOCUS
DEFINITION
ACCESSION
VERSION
KEYWORDS ORGANISM SOURCE

PAT 14-FEB-2001

linear

DNA

Unknown. Unclassified. REFERENCE AUTHORS TITLE JOURNAL FEATURES

Tobases 1 to 1308)
Sutliff, T.D. and Rodriguez, R.L.
Production of .alpha. .sub.1 -antitrypsin in plants
Patent: US 6127145-A 6 03-OCT-2000;
Location/Qualifiers

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                                                                                                                                                                                                                                                                                                                 236 GCTCGCGCACCAGTCCAACTCCACCAACATCTTCTTCAGCCCGGTGAGCATCGCCACCGC 295
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                                                                                                                                              335 GGAAGACCCTCAAGGCGACGCCGCTCAAAAACCGACACCAGTCATCACGACCAAGACCA 394
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                                                                             Length 1308;
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                                                                         Score 630.4; DB 6;
Pred. No. 2.1e-125;
0; Mismatches 346;
                             205 t
            /organism="unknown"
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70.88;
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                              1076 CGTGACGGAGGAGGCCCCCTGAAGCTCTCCAAGGCCGTGCACAAGGCGGTGCTCCACGAT 1135
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1295 TGTTACTGAAGAAGCTCCATTAAAATTGAGTAAAGCTGTTCACAAAGCCGTCTTAACTAT
                                                             1355 TGATGAAAAGGGTACCGAGGCCGCCGCGCTATGTTCCTGGAAGCTATTCCAATGAGCAT
                                                                                                                             1415 TCCACCAGAAGTTAAATTTAATAAACCATTCGTTTTTCTGATGATCGAGCAGAACACTAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                     1 (bases 1 to 1185)
Sutliff,T.D. and Rodriguez,R.L.
Production of alpha..sub.1 -antitrypsin in plants
Patent: US 6127145-A 3 03-ocr-2000;
                                                                                                                                                                                                                                                                                                         linear
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396 c 334 q
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/organism="Homo sapiens"
/db_xref="taxon:9606"
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1. .1345
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            1021 GACGAGAAGGGACGGACGCAGGGCCATGTTCCTGGAGGCCATCCCCATGTCCATC 1080
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                                                                                                                                                                                                                                                                          CAACATTTAGAGAATGAGTTGACTCATGACATTATTACTAAATTTTTAGAGAACGAGGAT 1175
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                                                                                                                                                                                GTTGAGAAAGGCACCCAGGGTAAGATCGTTGACCTAGTTAAAAGAATTAGATCGTGATACC
             GTCTTCGCACTAGTTAACTATATTTTTTCAAGGGTAAGTGGGAACGTCCTTTCGAGGTT
                                                                                                                         841 AGGCGCTCCGCTAGCCTCCACCTCCCGAAGCTGAGCATCACCGGCACGTACGACCTGAAG
                                                                               541 GTCTTCGCGCTCGTCAACTACATCTTCTTCAAGGGCAAGTGGGAGCGCCCGTTCGAGGTG
                                                                                                            AAAGATACTGAAGAGGAAGATTTTCATGTTGATCAAGTTACTACTGTCAAAGTTCCAATG
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M11465
Alpha-1-antitrypsin.
Human liver, cDNA to mRNA.
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/db_xref="cDB:G00-120-289"
//db_xref="cDB:G00-120-289"
//translation="MPSSVSWGILLAGLCCLVPVSLAEDPQGDAAQKTDTSHHDQDHP
TFNKITPNLAEFAFSLYRQLAHQSNSTNIFFSPVSIATAFAMLSLGTKADTHDEILEG
INFNLTEIPEAQIHEGFQELLRTLNQPDSGLQLTTGNGLFLSEGLKLVDKFLEDVKKL
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PDEGEKLQHLENELTHDITKTLENEDRRSABIHLPKLSITGYYDLKSVLGQLGITKV
FSNGADLSGVTERAPLKLSKAVKAVLTIDEKGTEAAGAMFLEAIPMSIPPEVKFNKP
FVFLMIEQNTKSPLFMGKVVNPTQK"
[2] sites; alpha-1-antitrypsin deficiency disease mutations. The Grante Falls (a one base pair deletion), Bellingham (a substitution) and Mattawa (a one base pair insertion) mutations all result in premature transcription termination.
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317 g 293 t
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Pred. No. 5.8e-85;
0; Mismatches 496;
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718 ACAAATTCCTAGAAGACGTCAAGAAACTATATCATAGTGAGGCTTTTACCGTTAATTTTG 777
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/organism="Homo sapiens"
/db_xref="taxon:9606"
1. 426 c 346 g 293
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                     14-MAR-1985 JP 1985051553
14-MAR-1984 US 84 58941
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                                                                                               strandedness: Both;
                                                                                                           topology: Linear;
hypothetical: No;
anti-sense: No;
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59.98;
JP 1986012289-A/1
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Maagaretsuto, W.I. and Guren, H.K.
REGION SPECIFIC MUTATION INDUCTION IN ALPHA-1-ANTI-TRYPSIN Patent: JP 1986012289-A 1 20-JAN-1986;
CHIMO JIEBETEITSUKUSU INC
                                                                                                                                                                                                                                                                                                                                                                                                        932 TGCCAGCTTACATTTACCCAAACTGTCCATTACTGGAACCTATGATCTGAAGAGCGTCCT
                                                                                                                                                                                                     452 CCTGAAGCTAGTGGATAAGTTTTTGGAGGATGTTAAAAAGTTGTACCACTCAGAAGCCTT
                                        164 TACCGTTAATTTTGGTGATACTGAGGAAGCTAAAAAGCAAATTAATGATTATGTTGAGAA
                                                       824 AGGCACCCAGGGTAAGATCGTTGACCTAGTTAAAGAATTAGATCGTGATACCGTCTTCGC
                                                                                                                                       884 ACTAGTTAACTATATTTTTTCAAGGGTAAGTGGGAACGTCCTTTCGAGGTTAAAGATAC
                                                                                                                                                     944 IGAAGAGGAAGATTTTCATGTTGATCAAGTTACTACTGTCAAAGTTCCAATGATGAAAAG
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                                                                                                                                                                                                                                               /product='alpha-1 antitrypsin' FT
.1193,'GCG')
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Pred. No. 5.1e-84;
AAACARETSUTO WAI INZUEII, GUREN HITOSHI KAWASAKI PC C12N15/00,A61K35/74,A61K37/64,C07H21/04,C07K15/04, PC C12P21/02/A61K37/04,
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                                                                                                                                                                                                                                                                                                   /note='amino acid 358=Lys'
replace(1142. .1144,'AAG')
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         'note='amino acid 342=Lys'
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Mon Dec

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linear PAT 21-MAY-1993
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838 AGAICGITGACCIAGITAAAGAATTAGAICGIGAIACCGICITCGCACIAGITAACIAIA 897
                                                                                                       TTTTTTTCAAGGGTAAGTGGGAACGTCCTTTCGAGGTTAAAGGTACTGAAGAGGAAGATT 957
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1 (bases 1 to 1312)

Kawasaski,G.H.
Glycolytic promotersfor regulated protein expression: protease inhibitor
                                                              561 GGACACCGAAGAGGCCAAGAAACAGATCAACGATTACGTGGAGAAGGGTACTCAAGGGA
                                                                                                                                                                                                                                                                  861 CCACCGCCATCTTCTTCCTGCTGATGAGGGGAAACTACAGCACCTGGAAAATGAACTCA
                                                                                                                                                                                                                                                                                                                                                                    Sequence 1 from Patent US 4599311.
102706
102706.1 GI:268359
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1547 - 16th Ave. East; Seattle, WA
Location/Qualifiers
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                                                                                                                                                                          358 CTCAAAAACCGACACCAGTCATCACGACCAAGACCATCCGACTTTTAATAAAATTACTC 417
                                                                                                                                                                                                                                   118 CAAATTTAGCCGAATTTGCTTTTTTTTTTTTAGACAATTAGCTCATCAAAGTAATTCTA 477
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                                                          Length 1312;
                                                       Score 436.4; DB 6; Length
Pred. No. 1.1e-83;
0; Mismatches 491; Indels
             281
/organism="unknown"
             324 g
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59.8%;
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1078 CTACTGCTATTTTTTTTTTACCAGAGGTAAGGTAAGCTTCAACATTTAGAGAATGAGTTGA 1137
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/db_xref="G1:15080499"
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59.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens, Similar to serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antirrypsin), member 1, clone MGC:9222 IMAGE:3859644, mRNA, complete cds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Center code: BCM-HGSC
Web site: http://www.hgsc.bcm.tmc.edu/cdna/
Contact: villalon@bcm.tmc.edu.
Villalon, D.K., Luna, R.A., Hale, S.M., Hulyk, S., Lu, X., Garcia,
Villalow, M., Telford, B, Hodgson, A., Bouck, J., Yu, W.,
Muzny, D.M., Gibbs, R.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 1584)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Submitted (30-JUL-2001) National Institutes of Health, Mammalian Gene Collection (MGC), Cancer Genomics Office, National Cancer Institute, 31 Center Drive, Room 11A03, Bethesda, MD 20892-2590,
1258 ITACCAAAGTTTTTTTTTAACGGTGCCGATTTGAGTGGTGTTACTGAAGAAGCTCCATTAA 1317
                        1318 AATTGAGTAAAGCTGTTCACAAAGCCGTCTTAACTATTGATGAAAAAGGGTACCGAGGCCG 1377
                                                                                                                                                      CCGGCGCTATGTTCCTGGAAGCTATTCCAATGAGCATTCCACCAGAAGTTAAATA 1437
                                                                                                                                                                                                                              1438 AACCATTCGTTTTTCTGATGATCGAGCAGAACACTAAAAGCCCCATTGTTATGGGTAAGG 1497
                                                                                                                                                                                                                                                      1082 AGCICICCAAGGCCGIGCAIAAGGCIGIGCIGACCAICGACGAGAAAGGGACIGAAGCIG 1141
                                                                                                                                                                                /codon_start=1
/product="Similar to serine (or cysteine) proteinase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cDNA Library Preparation: Life Technologies, Inc. cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Baylor College of Medicine Human Genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /tissue_type="Ovary, adenocarcinoma"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NIH-MGC Project URL: http://mgc.nci.nih.gov
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/db_xref="taxon:9606"
/clone="MGC:9222 IMAGE:3859644"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note="Vector: pCMV-SPORT6"
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/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: MGC help desk
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: DCTD/DTP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
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BC011991.1 GI:15080498
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Horrigan, S., Soppet, D.R. and Weaver, 2.
Cancer gene determination and therapeutic screening using signature
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                           1138 CTCATGACATTATTACTAAATTTTTAGAGAACGAGGATCGTCGTAGCGCTTCTCTGCACC 1197
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Location/Qualifiers
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349 a 386 c 325 g 292 t
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//translation="WESSVSWGILLIAGLCCLVPVSLAEDPOGDAAQKTDTSHHDQDH
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GLWFWITERPEAGLAGVGLAHQSTNIFFSPVSTATAFAMLSLGTKADTHDELLE
GLWFWITERPEAGLAGTALAAQPDSQLQLTGVAGLFLSGGLKLVDKFLEDVKK
LYHSEAFTVNFGDTEEAKKQINDXVEKGTGGKIVDLVKELDRDTYFALVNYIFFKGKW
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FLPDEGKLQHLENELTHDITTKFLENEDRRSASLHLPKLSITGTYDLKSVLGQLGITTK
VFSNGADLSGVTEEAPLKLSKAVHKAVLTIDEKGTEAAGAMFLEAIPMSIPPEVKFNK
PFVFLMIEQNTKSPLEMGKVVNPTQK"
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      PRI 01-NOV-1994
                                                                                                                                             1 (bases I to 1231)
Bollen, A., Herzog, A., Cravador, A., Herion, P., Chuchana, P., Vander Straten, A., Loriau, R., Jacobs, P. and van Elsen, A. Conjug and expression in Escherichia coli of full-length Complementary DNA coding for human alpha 1-antitrypsin 84107980
                                                                                                                                                                                                                                                                                                                                                                                   Revised sequence of full-length complementary DNA coding for human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Courtney, M., Buchwalder, A., Tessier, L.H., Jaye, M., Benavente, A., Balland, A., Kohli, V., Lathe, R., Tolstoshev, P. and Lecocq, J.P. High-level production of biologically active human alpha 1-antitrypsin in Escherichia coli and Proc. Natl. Acad. Sci. U.S.A. 81 (3), 669-673 (1984)
                                                                                       alpha-1-antitrypsin; antiprotease; antitrypsin.
Human liver, cDNA to mRNA, clones pULB1523 [1] and pTG603 [3]
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          Human alpha-1-antitrypsin mRNA, complete cds.
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/db_xref="GI:177829"
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Best Local Similarity 59.7%;
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/citation=[1]
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Young, P.E., Augustus, M., Carter, K.C., Ebner, R., Endress, G., Horrigan, S., Soppet, D.R. and Weaver, Z. Cancer gene determination and therapeutic screening using signature
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Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                  CTACTGCTATTTTTTTTACCAGACGAAGGTAAGCTTCAACATTTAGAGAATGAGTTGA 11.37
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Sequence 5847 from Patent WO0194629.
AX335338 GI:18126057
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Score 433.2; DB 6; Pred. No. 5.5e-83; 0; Mismatches 493;

28.4%;

Best Local Similarity 59.7 Matches 729; Conservative

Query Match

Length 1371;

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1093 AGCTCTCCAAGGCCGTGCATAAGGCTGTGCTGACCATCGACGAGAAGGGACTGAAGCTG 1152
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                                                                    CTCAAAAAACGACACCAGTCATCACGACCAAGACCATCGGACTTTTAATAAAATTACTC 417
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PTRKTTPNTATPREAFSLYFQLAHQSVSTNIFFSPSYSTATFATADTHDELLE
GLRNLTEIPEAQHTEGFQELLRTLMQPDSQLQLTFGNGLELSEGLKLVDKFLEDVK
LYHSEAFTVNFGDTEEAKQINDYVEKGTGGKIVDLVKELDBDYVFALVWYTVRYPWRKLAGWTNOGKKKELSSWLLAWYLFKTGWRF
PERPEWKOTPEEDFHVDQYTVWYRPWRKRAGWHVDGKKKESWLLAWYLGKYTGRATAIF
FLPDEGKLGHLENETHDITTFFLENEDRESALHLPKLSTGTYDLKSVLGGLGTTK
VFSNGADLSGVTEEAPLKLSKAVHKAVLTIDEKGTEAAGAMFLEAIPMSIPPEVKFNK
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Colau,B., Chuchana,P. and Bollen,A.
Revised sequence of full-length complementary DNA coding for human
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                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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1378 CCGGCGCTAIGTICCTGGAAGCTATTCCAATGAGCATTCCACCAGAAGTTAAATTTAATA 1437
                                     1153 CTGGGGCCATGTTTTAGAGGCCATACCCATGTCTATCCCCCCCGGGGTCAAGTTCAACA 1212
                                                                                         1 (bases 1 to 274)

Kurachi, K., Chandra, T., Degen, S.J., White, T.T., Marchioro, T.L., Woo, S.L. and Davie, E.W.
Woo, S.L. and sequence of cDNA coding for alpha 1-antitrypsin Proc. Natl. Acad. Sci. U.S.A. 78 (11), 6826-6830 (1981)
82082539
                                                                        AACCATTCGTTTTTCTGATGATCGAGCAGAACACTAAAAGCCCATTGTTTATGGGTAAGG
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Location/Qualifiers
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/protein_id="CAA28838.1"
/db_xref="CI:28966"
/db_xref="SMISS-PROT:P01009"
                                                                                                                                                                                                                                                                               mRNA
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Nature 312 (5989), 77-80 (1984)
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                                                                                                                                              Length 1371;
                                                                                                                                                                                     Indels
                                                                                                                                                  Score 433.2; DB 9;
Pred. No. 5.5e-83;
0; Mismatches 493;
                                                                                                                                                DB 9;
PFVFLMIEQNTKSPLFMGKVVNPTQK"
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                     111. .1292
/product="unnamed"
1371
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59.7%;
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                                                                                                                                                             Best Local Similaricy
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111. .129
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PRI 29-SEP-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Submitted (29-AUG-2000) Sumio Sugano, Institute of Medical Science, University of Tokyo, Laboratory of Genome Structure Analysis, Human Genome Center; Shirokane-dai, 4-6-1, Minato-ku, Tokyo 108-8639, Japan (E-mail:canal@ims.u-tokyo.ac.jp, Tel:81-3-5449-5286,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ဌ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kawabata, A., Hikiji, T., Kobatake, N., Inagaki, H., Ikema, Y., Okamoto, S., Okitani, R., Ota, T., Suzuki, Y., Obayashi, M., Nishi, T., Shibahara, T., Tanaka, T., Nakamura, Y., Isogai, T. and Sugano, S. M. Debo human cDNA sequencing project
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NEDO human CDNA sequencing project supported by Ministry of International Trade and Industry of Japan; CDNA full insert sequencing: Research Association for Blotechnology: CDNA library construction, 5'- & 3'-end one pass sequencing: Departent of Virology and Human Genome Center, Institute of Medical Science, University of Tokyo (partly supported by Science and Technology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ARO26174 1139 bp mRNA 11near PRI 29-5EP-2,
Homo sapiens cDNA: FLJ22521 fis, clone HRC12386, highly similar
AF113676 Homo sapiens clone FLB2803 PRO0684 mRNA.
1033 rcacraaggrerreaggaarggggergacerecegggreacagagggggegecerea 1092
                                                               1258 TTACCAAAGTTTTTTTTAACGGTGCCGATTTGAGTGGTGTTACTGAAGAAGCTCCATTAA 1317
                                                                                                                                                         AATTGAGTAAAGCTGTTCACAAAGCCGTCTTAACTATTGATGAAAAGGGTACCGAGGCCG 1377
                                                                                                                                                                                                                                                 CCGGCGCTATGTTCCTGGAAGCTATTCCAATGAGCATTCCACCAGAAGTTAAATTAATA 1437
                                                                                                                                                                                                                                                                                         CIGGGGCCATGITITAGAGGCCATACCCATGICIATCCCCCCGAGGICAAGITCAACA 1212
                                                                                                                                                                                                                                                                                                                                                                oligo capping; fis (full insert sequence).
Homo sapiens primary human renal epithelial cells cDNA to mRNA,
clone_lib:HRC clone:HRC12386.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sugano,S., Suzuki,Y., Ota,T., Obayashi,M., Nishi,T., Isogai,T., Shibahara,T., Tanaka,T. and Nakamura,Y.
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/cell_type="primary human renal epithelial cells"
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/note="cloning vector pME18SFL3"
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/db_xref="taxon:9606"
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AK026174
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KEYWORDS
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Length 1399;
                                         Indels
                                       0; Mismatches 493;
    DB 9;
Score 433.2; DB 9
Pred. No. 5.5e-83;
28.4%;
59.7%;
                                       Matches 729; Conservative
                    Best Local Similarity
  Query Match
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PN JP1984091886-A/1
PD 26-MAY-1984
PF 12-MG-1983 JP 1983147907
PF 13-AG-1983 US 83 489406 PI
PR 13-AG-1982 US 82 408099, 28-APR-1983 US 83 489406 PI
PG 13-AG-1982 US 82 408099, 28-APR-1983 US 83 489406 PI
PC 21015/00, A61K35/72, A61K37/64, CO7767/00, CO7721/04, C12N1/16, PC
C12N9/99,
                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                        1158 CTGGGGCCATGTTTTAGAGGCCATACCCATGTCTATCCCCCCGGGGTCAAGTTCAACA 1217
                                                                                                    AACCATTCGTTTTTCTGATGATCGAGCAGAACACTAAAAGCCCATTGTTTATGGGTAAGG 1497
                                                                                                                     AGCTCTCCAAGGCCGTGCATAAGGCTGTGCTGACCATCGACGAGAAAGGGACTGAAGCTG 1157
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                                                                                                                                                                                                                                                                                                                                                                                                1 (bases 1 to 1434)
Guren,E.K. and Richiyaado,U.
SUGAR LYSING PROMOTOR : PROTEASE INHIBITOR FOR DEVELOPMENT
                                              CCGGCGCTATGTTCCTGGAAGCTATTCCAATGAGCATTCCACCAGAAGTTAAATA
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3S 47..1303
UTR 1304
1..1434
1..1434
47..118
47..118
1L_peptide 119..1300
Location/Qualifiers
                                                                                                                                                                                                                                                                  linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 433.2; DB 6
Pred. No. 5.5e-83;
                                                                                                                                                                                                                                                                   RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Patent: JP 1984091886-A 1 26-MAY-1984;
JIMOSU CORP
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strandedness: Double;
topology: Linear;
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/organism="Homo sapiens"
/db_xref="taxon:9606"
432 c 340 g 29:
                                                                                                                                                                                                                                                            1434 bp cDNA encoding human antitrypsin. E00195. E0195. E0195. B1984091886-A/1. Homo saviora
                                                                                                                                                                TIGICAACCCAACTCAGAAGTA 1519
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anti-sense: No;
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CAAATTTAGCCGAATTTGCTTTTTTTTTTGTATAGACAATTAGCTCATCAAAGTAATTCTA 477
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therapeutit formulations, and vectors and bacteria for such method
and their production
Patent: EP 0137633-Al 3 17-APR-1985;
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                                                                                                                                                                                                                                                                                                                  293
                                                                                   Sequence 3 from Patent EP 0137633.
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Parker, M.L. and Kawasaki, G.H.
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TTGTCAACCCAACTCAGAAGTA 1519
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PAT 02-DEC-1994
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Russell.P.R.
Method for expressing foreign genes in schizosaccharomyces pombe and the use in therapeutic formulations of the products, DNA constructs and transformant strains of schizosaccharomyces pombe usable in such method and their preparation
patent: EP 0139383-A1 1 02-MAY-1985;
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                                                                   201 CCAACCTGGCTGAGTTCGCCTTCAGCCTATACCGCCAGCTGGCACACCAGTCCAACAGCA
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Kawasaki, G.H. and Woodbury, R.G.
Glycolytic promoters for regulated protein expression: protease
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